PCT/IL2004/001049

FIGURE 1A.

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5	ATGTCACTGA	AAAACGAGCC	ACGGGTAAAT	ACCTCTGCAC	TGCAGAAAAT	TGCTGCTGAC	ATGAGTAATA	70
J	TCATAGAAAA	TCTGGACACG	CGGGAACTCC	ACTTTGAGGG	AGAGGAGGTA	GACTACGACG	TGTCTCCCAG	140
	CGATCCCAAG	ATACAAGAAG	TGTATATCCC	TTTCTCTGCT	ATTTATAACA	CTCAAGGATT	TAAGGAGCCT	210
1 0	AATATACAGA	CGTATCTCTC	CGGCTGTCCA	ATAAAAGCAC	AAGTTCTGGA	AGTGGAACGC	TTCACATCTA	280
	CAACAAGGGT	ACCAAGTATT	AATCTTTACA	CTATTGAATT	AACACATGGG	GAATTTAAAT	GGCAAGTTAA	350
15	GAGGAAATTC	AAGCATTTTC	AAGAATTTCA	CAGAGAGCTG	CTCAAGTACA	AAGCCTTTAT	CCGCATCCCC	420
13	ATTCCCACTA	GAAGACACAC	GTTTAGGAGG	CAAAACGTCA	GAGAGGAGCC	TCGAGAGATG	CCCAGTTTGC	490
	CCCGTTCATC	TGAAAACATG	ATAAGAGAAG	AACAATTCCT	TGGTAGAAGA	AAACAACTGG	AAGATTACTT	560
20	GACAAAGATA	CTAAAAATGC	CCATGTATAG	AAACTATCAT	GCCACAACAG	AGTTTCTTGA	TATAAGCCAG	630
	CTGTCTTTCA	TCCATGATTT	GGGACCAAAG	GGCATAGAAG	GTATGATAAT	GAAAAGATCT	GGAGGACACA	700
25	GAATACCAGG	CTTGAATTGC	TGTGGTCAGG	GAAGAGCCTG	CTACAGATGG	TCAAAAAGAT	GGTTAATAGT	770
23	GAAAGATTCC	TTTTTATTGT	ATATGAAACC	AGACAGCGGT	GCCATTGCCT	TCGTCCTGCT	GGTAGACAAA	840
	GAATTCAAAA	TTAAGGTGGG	GAAGAAGGAG	ACAGAAACGA	AATATGGAAT	CCGAATTGAT	AATCTTTCAA	910
30	GGACACTTAT	TTTAAAATGC	AACAGCTATA	GACATGCTCG	GTGGTGGGGA	GGGGCTATAG	AAGAATTCAT	980
	CCAGAAACAT	GGCACCAACT	TTCTCAAAGA	TCATCGATTT	GGGTCATATG	CTGCTATCCA	AGAGAATGCT	1050
35	TTAGCTAAAT	GGTATGTTAA	TGCCAAAGGA	TATTTTGAAG	ATGTGGCAAA	TGCAATGGAA	GAGGCAAATG	1120
55	AAGAGATTTT	TATCACAGAC	TGGTGGCTGA	GTCCAGAAAT	CTTCCTGAAA	CGCCCAGTGG	TTGAGGGAAA	1190
	TCGTTGGAGG	TTGGACTGCA	TTCTTAAACG	AAAAGCACAA	CAAGGAGTGA	GGATCTTCAT	AATGCTCTAC	1260
40	AAAGAGGTGG	AACTCGCTCT	TGGCATCAAT	AGTGAATACA	CCAAGAGGAC	TTTGATGCGT	CTACATCCCA	1330
	ACATAAAGGT	GATGAGACAC	CCGGATCATG	TGTCATCCAC	CGTCTATTTG	TGGGCTCACC	ATGAGAAGCT	1400
45	TGTCATCATT	GACCAATCGG	TGGCCTTTGT	GGGAGGGATT	GACCTGGCCT	ATGGAAGGTG	GGACGACAAT	1470
	GAGCACAGAC	TCACAGACGT	GGGCAGTGTG	AAGCGGGTCA	CTTCAGGACC	GTCTCTGGGT	TCCCTCCCAC	1540
	CTGCCGCAAT	GGAGTCTATG	GAATCCTTAA	GACTCAAAGA	TAAAAATGAG	CCTGTTCAAA	ACCTACCCAT	1610
50	CCAGAAGAGT	ATTGATGATG	TGGATTCAAA	ACTGAAAGGA	ATAGGAAAGC	CAAGAAAGTT	CTCCAAATTT	1680
	AGTCTCTACA	AGCAGCTCCA	CAGGCACCAC	CTGCACGACG	CAGATAGCAT	CAGCAGCATT	GACAGCACCT	1750
55	CCAGTTATTT	TAATCACTAT	AGAAGTCATC	ACAATTTAAT	CCATGGTTTA	AAACCCCACT	TCAAACTCTT	1820
	TCACCCGTCC	AGTGAGTCTG	AGCAAGGACT	CACTAGACCT	CATGCTGATA	CCGGGTCCAT	CCGTAGTTTA	1890
	CAGACAGGTG	TGGGAGAGCT	GCATGGGGAA	ACCAGATTCT	GGCATGGAAA	GGACTACTGC	AATTTCGTCT	1960
60	TCAAAGACTG	GGTTCAACTT	GATAAACCTT	TTGCTGATTT	CATTGACAGG	TACTCCACGC	CCCGGATGCC	2030
	CTGGCATGAC	ATTGCCTCTG	CAGTCCACGG	GAAGGCGGCT	CGTGATGTGG	CACGTCACTT	CATCCAGCGC	2100
65	TGGAACTTCA	CAAAAATTAT	GAAATCAAAA	TATCGGTCCC	TTTCTTATCC	TTTTCTGCTT	CCAAAGTCTC	2170
	AAACAACAGC	CCATGAGTTG	AGATATCAAG	TGCCTGGGTC	TGTCCATGCT	AACGTACAGT	TGCTCCGCTC	2240
	TGCTGCTGAT	TGGTCTGCTG	GTATAAAGTA	CCATGAAGAG	TCCATCCACG	CCGCTTACGT	CCATGTGATA	2310
70	GAGAACAGCA	GGCACTATAT	CTATATCGAA	AACCAGTTTT	TCATAAGCTG	TGCTGATGAC	AAAGTTGTGT	2380

FIGURE 1B.

5	TCAACAAGAT	AGGCGATGCC	ATTGCCCAGA	GGATCCTGAA	AGCTCACAGG	GAAAACCAGA	AATACCGGGT	245
,	ATATGTCGTG	ATACCACTTC	TGCCAGGGTT	CGAAGGAGAC	ATTTCAACCG	GCGGAGGAAA	TGCTCTACAG	252
	GCAATCATGC	ACTTCAACTA	CAGAACCATG	TGCAGAGGAG	AAAATTCCAT	CCTTGGACAG	TTAAAAGCAG	259
10	AGCTTGGTAA	TCAGTGGATA	AATTACATAT	CATTCTGTGG	TCTTAGAACA	CATGCAGAGC	TCGAAGGAAA	266
	CCTAGTAACT	GAGCTTATCT	ATGTCCACAG	CAAGTTGTTA	ATTGCTGATG	ATAACACTGT	TATTATTGGC	273
15	TCTGCCAACA	TAAATGACCG	CAGCATGCTG	GGAAAGCGTG	ACAGTGAAAT	GGCTGTCATT	GTGCAAGATA	280
J	CAGAGACTGT	TCCTTCAGTA	ATGGATGGAA	AAGAGTACCA	AGCTGGCCGG	TTTGCCCGAG	GACTTCGGCT	287
	ACAGTGCTTT	AGGGTTGTCC	TTGGCTATCT	TGATGACCCA	AGTGAGGACA	TTCAGGATCC	AGTGAGTGAC	294
20	AAATTCTTCA	AGGAGGTGTG	GGTTTCAACA	GCAGCTCGAA	ATGCTACAAT	TTATGACAAG	GTTTTCCGGT	301
	GCCTTCCCAA	TGATGAAGTA	CACAATTTAA	TTCAGCTGAG	AGACTTTATA	AACAAGCCCG	TATTAGCTAA	3080
2.5	GGAAGATCCC	ATTCGAGCTG	AGGAGGAACT	GAAGAAGATC	CGTGGATTTT	TGGTGCAATT	CCCCTTTTAT	3150
	TTCTTGTCTG	AAGAAAGCCT	ACTGCCTTCT	GTTGGGACCA	AAGAGGCCAT	AGTGCCCATG	GAGGTTTGGA	3220
	CTTAA 3225							

FIGURE 2.

5								
J	MSLKNEPRVN	TSALQKIAAD	MSNIIENLDT	RELHFEGEEV	DYDVSPSDPK	IQEVYIPFSA	IYNTQGFKEP	70
	NIQTYLSGCP	1KAQVLEVER	FTSTTRVPSI	NLYTIELTHG	EFKWQVKRKF	KHFQEFHREL	LKYKAFIRIP	140
10	IPTRRHTFRR	QNVREEPREM	PSLPRSSENM	IREEQFLGRR	KQLEDYLTKI	LKMPMYRNYH	ATTEFLDISQ	210
	LSFIHDLGPK	GIEGMIMKRS	GGHRIPGLNC	CGQGRACYRW	SKRWLIVKDS	FLLYMKPDSG	AIAFVLLVDK	280
15	EFKIKVGKKE	TETKYGIRID	NLSRTLILKC	NSYRHARWWG	GAIEEFIQKH	GTNFLKDHRF	GSYAAIQENA	350
13	LAKWYVNAKG	YFEDVANAME	EANEEIFITD	WWLSPEIFLK	RPVVEGNRWR	LDCILKRKAQ	QGVRIFIMLY	420
	KEVELALGIN	SEYTKRTLMR	LHPNIKVMRH	PDHVSSTVYL	WAHHEKLVII	DQSVAFVGGI	DLAYGRWDDN	490
20	EHRLTDVGSV	KRVTSGPSLG	SLPPAAMESM	ESLRLKDKNE	PVQNLPIQKS	IDDVDSKLKG	IGKPRKFSKF	560
	slykQlhrhh	LHDADSISSI	DSTSSYFNHY	RSHHNLIHGL	KPHFKLFHPS	SESEQGLTRP	HADTGSIRSL	630
25	QTGVGELHGE	TRFWHGKDYC	NEAEKDMAÓT	DKPFADFIDR	YSTPRMPWHD	IASAVHGKAA	RDVARHFIQR	700
23	WNFTKIMKSK	YRSLSYPFLL	PKSQTTAHEL	RYQVPGSVHA	nvqllrsaad	WSAGIKYHEE	SIHAAYVHVI	770
	ENSRHYIYIE	NQFFISCADD	KVVFNKIGDA	IAQRILKAHR	ENQKYRVYVV	IPLLPGFEGD	ISTGGGNALQ	840
30	AIMHFNYRTM	CRGENSILGQ	LKAELGNQWI	NYISFCGLRT	HAELEGNLVT	ELIYVHSKLL	IADDNTVIIG	910
	SANINDRSML	GKRDSEMAVI	VQDTETVPSV	MDGKEYQAGR	FARGLRLQCF	RVVLGYLDDP	SEDIQDPVSD	980
35	KFFKEVWVST	AARNATIYDK	VFRCLPNDEV	HNLIQLRDFI	NKPVLAKEDP	IRAEEELKKI	RGFLVQFPFY	1050
,,	FLSEESLLPS	VGTKEAIVPM	EVWT 1074					

FIGURE 3A.

	ATGACGGCGA	CCCCTGAGAG	CCTCTTCCCC	ACTGGGGACG	AACTGGACTC	CAGCCAGCTC	CAGATGGAGT	70
5	CCGATGAGGT	GGACACCCTG	AAGGAGGGAG	AGGACCCAGC	CGACCGGATG	CACCCGTTTC	TGGCCATCTA	140
	TGAGCTTCAG	TCTCTGAAAG	TGCACCCCTT	GGTGTTCGCA	CCTGGGGTCC	CTGTCACAGC	CCAGGTGGTG	210
10	GGCACCGAAA	GATATACCAG	CGGATCCAAG	GTGGGAACCT	GCACTCTGTA	TTCTGTCCGC	TTGACTCACG	280
10	GCGACTTTTC	CTGGACAACC	AAGAAGAAAT	ACCGTCATTT	TCAGGAGCTG	CATCGGGACC	TCCTGAGACA	350
	CAAAGTCTTG	ATGAGTCTGC	TCCCTCTGGC	TCGATTTGCC	GTTGCCTATT	CTCCAGCCCG	AGATGCAGGC	420
15	AACAGAGAGA	TGCCCTCTCT	ACCCCGGGCA	GGTCCTGAGG	GCTCCACCAG	ACATGCAGCC	AGCAAACAGA	490
	AATACCTGGA	GAATTACCTC	AACTGTCTCT	TGACCATGTC	TTTCTATCGC	AACTACCATG	CCATGACAGA	560
20	GTTCCTGGAA	GTCAGTCAGC	TGTCCTTTAT	CCCGGACTTG	GGCCGCAAAG	GACTGGAGGG	GATGATCCGG	630
20	AAGCGCTCAG	GTGGCCACCG	TGTTCCTGGC	CTCACCTGCT	GTGGCCGAGA	CCAAGTTTGT	TATCGCTGGT	700
	CCAAGAGGTG	GCTGGTGGTG	AAGGACTCCT	TCCTGCTGTA	CATGTGCCTC	GAGACAGGTG	CCATCTCATT	770
25	TGTTCAGCTC	TTTGACCCTG	GCTTTGAGGT	GCAAGTGGGG	AAAAGGAGCA	CGGAGGCACG	GCACGGCGTG	840
	CGGATCGATA	CCTCCCACAG	GTCCTTGATT	CTCAAGTGCA	GCAGCTACCG	GCAGGCACGG	TGGTGGGCCC	910
30	AAGAGATCAC	TGAGCTGGCA	CAGGGCCCAG	GCAGAGACTT	CCTACAGCTG	CACCGGCATG	ACAGCTACGC	980
50	CCCACCCCGG	CCTGGGACCT	TGGCCCGGTG	GTTTGTGAAT	GGGGCAGGTT	ACTTTGCTGC	TGTGGCAGAT	1050
	GCCATCCTTC	GAGCTCAAGA	GGAGATTTTC	ATCACAGACT	GGTGGTTGAG	TCCTGAGGTT	TACCTGAAGC	1120
35	GTCCGGCCCA	TTCAGATGAC	TGGAGACTGG	ACATTATGCT	CAAGAGGAAG	GCGGAGGAAG	GTGTCCGTGT	1190
	GTCTATTCTG	CTGTTTAAAG	AAGTGGAATT	GGCCTTGGGC	ATCAACAGTG	GCTATAGCAA	GAGGGCGCTG	1260
40	ATGCTGCTGC	ACCCCAACAT	AAAGGTGATG	CGTCACCCAG	ACCAAGTGAC	GTTGTGGGCC	CATCATGAGA	1330
-10	AGCTCCTGGT	GGTGGACCAA	GTGGTAGCAT	TCCTGGGGGG	ACTGGACCTT	GCCTATGGCC	GCTGGGATGA	1400
	CCTGCACTAC	CGACTGACTG	ACCTTGGAGA	CTCCTCTGAA	TCAGCTGCCT	CCCAGCCTCC	CACCCCGCGC	1470
45	CCAGACTCAC	CAGCCACCCC	AGACCTCTCT	CACAACCAAT	TCTTCTGGCT	GGGCAAGGAC	TACAGCAATC	1540
	TTATCACCAA	GGACTGGGTG	CAGCTGGACC	GGCCTTTCGA	AGATTTCATT	GACAGGGAGA	CGACCCCTCG	1610
50	GATGCCATGG	CGGGACGTTG	GGGTGGTCGT	CCATGGCCTA	CCGGCCCGGG	ACCTTGCCCG	GCACTTCATC	1680
20	CAGCGCTGGA	ACTTCACCAA	GACCACCAAG	GCCAAGTACA	AGACTCCCAT	ATACCCCTAC	CTGCTTCCCA	1750
•	AGTCTACCAG	CACGGCCAAT	CAGCTCCCCT	TCACACTTCC	AGGAGGCAG	TGCACCACCG	TACAGGTCTT	1820
55	GCGATCAGTG	GACCGCTGGT	CAGCAGGGAC	TCTGGAGAAC	TCCATCCTCA	ATGCCTACCT	GCACACCATC	1890
	AGGGAGAGCC	AGCACTTCCT	CTACATTGAG	AATCAGTTCT	TCATTAGCTG	CTCAGATGGG	CGGACGGTTC	1960
60	TGAACAAGGT	GGGCGATGAG	ATTGTGGACA	GAATCCTGAA	GGCCCACAAA	CAGGGGTGGT	GTTACCGAGT	2030
00	CTACGTGCTT	TTGCCCTTAC	TCCCTGGCTT	CGAGGGTGAC	ATCTCCACGG	GCGGTGGCAA	CTCCATCCAG	2100
•	GCCATTCTGC	ACTTTACTTA	CAGGACCCTG	TGTCGTGGGG	AGTATTCAAT	CCTGCATCGC"	CTTAAAGCAG	2170
65	CCATGGGGAC	AGCATGGCGG	GACTATATTT	CCATCTGCGG	GCTTCGTACA	CACGGAGAGC	TGGGCGGGCA	2240
	CCCCGTCTCG	GAGCTCATCT	ACATCCACAG	CAAGGTGCTC	ATCGCAGATG	ACCGGACAGT	CATCATTGGT	2310
70	TCTGCAAACA	TCAATGACCG	GAGCTTGCTG	GGGAAGCGGG	ACAGTGAGCT	GGCCGTGCTG	ATCGAGGACA	2380
, 5	CAGAGACGGA	ACCATCCCTC	ATGAATGGGG	CAGAGTATCA	GGCGGGCAGG	TTTGCCTTGA	GTCTGCGGAA	2450
	GCACTGCTTC	GGTGTGATTC	TTGGAGCAAA	TACCCGGCCA	GACTTGGATC	TCCGAGACCC	CATCTGTGAT	2520

FIGURE 3B.

5 GACTTCTTCC AGTTGTGGCA AGACATGGCT GAGAGCAACG CCAATATCTA TGAGCAGATC TTCCGCTGCC 2590
TGCCATCCAA TGCCACGCGT TCCCTGCGGA CTCTCCGGGA GTACGTGGCC GTGGAGCCCT TGGCCACGGT 2660
CAGTCCCCCC TTGGCTCGGT CTGAGCTCAC CCAGGTCCAG GGCCACCTGG TCCACTTCCC CCTCAAGTTC 2730
10 CTAGAGGATG AGTCTTTGCT GCCCCGCTG GGTAGCAAGG AGGGCATGAT CCCCCTAGAA GTGTGGACAT 2800
AG 2802

FIGURE 4.

	3	MTATPESLFP	TGDELDSSQL	QMESDEVDTL	KEGEDPADRM	HPFLAIYELQ	SLKVHPLVFA	PGVPVTAQVV	70
		GTERYTSGSK	VGTCTLYSVR	LTHGDFSWTT	KKKYRHFQEL	HRDLLRHKVL	MSLLPLARFA	VAYSPARDAG	140
	10	NREMPSLPRA	GPEGSTRHAA	SKQKYLENYL	NCLLTMSFYR	NYHAMTEFLE	VSQLSFIPDL	GRKGLEGMIR	210
	10	KRSGGHRVPG	LTCCGRDQVC	YRWSKRWLVV	KDSFLLYMCL	ETGAISFVQL	FDPGFEVQVG	KRSTEARHGV	280
		RIDTSHRSLI	LKCSSYRQAR	WWAQEITELA	QGPGRDFLQL	HRHDSYAPPR	PGTLARWFVN	GAGYFAAVAD	350
	15	AILRAQEEIF	ITDWWLSPEV	YLKRPAHSDD	WRLDIMLKRK	AEEGVRVSIL	LFKEVELALG	INSGYSKRAL	420
		MLLHPNIKVM	RHPDQVTLWA	HHEKLLVVDQ	VVAFLGGLDL	AYGRWDDLHY	RLTDLGDSSE	SAASQPPTPR	490
20	20	PDSPATPDLS	HNQFFWLGKD	YSNLITKDWV	QLDRPFEDFI	DRETTPRMPW	RDVGVVVHGL	PARDLARHFI	560
	20	QRWNFTKTTK	AKYKTPIYPY	LLPKSTSTAN	QLPFTLPGGQ	CTTVQVLRSV	DRWSAGTLEN	SILNAYLHTI	630
		RESQHFLYIE	NQFFISCSDG	RTVLNKVGDE	IVDRILKAHK	QGWCYRVYVL	LPLLPGFEGD	ISTGGGNSIQ	700
	25	AILHFTYRTL	CRGEYSILHR	LKAAMGTAWR	DYISICGLRT	HGELGGHPVS	ELIYIHSKVL	IADDRTVIIG	770
		SANINDRSLL	GKRDSELAVL	IEDTETEPSL	MNGAEYQAGR	FALSLRKHCF	GVILGANTRP	DLDLRDPICD	840
	30	DFFQLWQDMA	ESNANIYEQI	FRCLPSNATR	SLRTLREYVA	VEPLATVSPP	LARSELTQVQ	GHLVHFPLKF	910
	20	LEDESLLPPL	GSKEGMIPLE	VWT 933					

FIGURE 5.

ATGAAGCCTA AACTGATGTA CCAGGAGCTG AAGGTGCCTG CAGAGGAGCC CGCCAATGAG CTGCCCATGA 70 ATGAGATTGA GGCGTGGAAG GCTGCGGAAA AGAAAGCCCG CTGGGTCCTG CTGGTCCTCA TTCTGGCGGT 140 5 TGTGGGCTTC GGAGCCCTGA TGACTCAGCT GTTTCTATGG GAATACGGCG ACTTGCATCT CTTTGGGCCC 210 AACCAGCGCC CAGCCCCCTG CTATGACCCT TGCGAAGCAG TGCTGGTGGA AAGCATTCCT GAGGGCCTGG 280 ACTTCCCCAA TGCCTCCACG GGGAACCCTT CCACCAGCCA GGCCTGGCTG GGCCTGCTCG CCGGTGCGCA 350 10 CAGCAGCCTG GACATCGCCT CCTTCTACTG GACCCTCACC AACAATGACA CCCACACGCA GGAGCCCTCT 420 GCCCAGCAGG GTGAGGAGGT CCTCCGGCAG CTGCAGACCC TGGCACCAAA GGGCGTGAAC GTCCGCATCG 490 15 CTGTGAGCAA GCCCAGCGGG CCCCAGCCAC AGGCGGACCT GCAGGCTCTG CTGCAGAGCG GTGCCCAGGT 560 CCGCATGGTG GACATGCAGA AGCTGACCCA TGGCGTCCTG CATACCAAGT TCTGGGTGGT GGACCAGACC 630 CACTTCTACC TGGGCAGTGC CAACATGGAC TGGCGTTCAC TGACCCAGGT CAAGGAGCTG GGCGTGGTCA 700 20 TGTACAACTG CAGCTGCCTG GCTCGAGACC TGACCAAGAT CTTTGAGGCC TACTGGTTCC TGGGCCAGGC 770 AGGCAGCTCC ATCCCATCAA CTTGGCCCCG GTTCTATGAC ACCGCTACA ACCAAGAGAC ACCAATGGAG 840 25 ATCTGCCTCA ATGGAACCCC TGCTCTGGCC TACCTGGCGA GTGCGCCCCC ACCCCTGTGT CCAAGTGGCC 910 GCACTCCAGA CCTGAAGGCT CTACTCAACG TGGTGGACAA TGCCCGGAGT TTCATCTACG TCGCTGTCAT 980 30 GAACTACCTG CCCACTCTGG AGTTCTCCCA CCCTCACAGG TTCTGGCCTG CCATTGACGA TGGGCTGCGG 1050 CGGGCCACCT ACGAGCGTGG CGTCAAGGTG CGCCTGCTCA TCAGCTGCTG GGGACACTCG GAGCCATCCA 1120 TGCGGGCCTT CCTGCTCTCT CTGGCTGCCC TGCGTGACAA CCATACCCAC TCTGACATCC AGGTGAAACT 1190 35 CTTTGTGGTC CCCGCGGATG AGGCCCAGGC TCGAATCCCA TATGCCCGTG TCAACCACAA CAAGTACATG 1260 GTGACTGAAC GCGCCACCTA CATCGGAACC TCCAACTGGT CTGGCAACTA CTTCACGGAG ACGGCGGGCA 1330 CCTCGCTGCT GGTGACGCAG AATGGGAGGG GCGGCCTGCG GAGCCAGCTG GAGGCCATTT TCCTGAGGGA 1400 40 CTGGGACTCC CCTTACAGCC ATGACCTTGA CACCTCAGCT GACAGCGTGG GCAACGCCTG CCGCCTGCTC 1470 TGA 1473 45

50

55

FIGURE 6.

	MKPKLMYQEL	KVPAEEPANE	LPMNEIEAWK	AAEKKARWVL	LVLILAVVGF	GALMTQLFLW	60
	EYGDLHLFGP	NQRPAPCYDP	CEAVLVESIP	EGLDFPNAST	GNPSTSQAWL	GLLAGAHSSL	120
5	DIASFYWTLT	NNDTHTQEPS	AQQGEEVLRQ	LQTLAPKGVN	VRIAVSKPSG	PQPQADLQAL	180
	LQSGAQVRMV	DMQKLTHGVL	HTKFWVVDQT	HFYLGSANMD	WRSLTQVKEL	GVVMYNCSCL	240
	ARDLTKIFEA	YWFLGQAGSS	IPSTWPRFYD	TRYNQETPME	ICLNGTPALA	YLASAPPPLC	300
	PSGRTPDLKA	LLNVVDNARS	FIYVAVMNYL	PTLEFSHPHR	FWPAIDDGLR	RATYERGVKV	360
,	RLLISCWGHS	EPSMRAFLLS	LAALRONHTH	SDIQVKLFVV	PADEAQARIP	YARVNHNKYM	420
10	VTERATYIGT	SNWSGNYFTE	TAGTSLLVTQ	NGRGGLRSQL	EAIFLRDWDS	PYSHDLDTSA	480
	DSVGNACRLL	490					

FIGURE 7.

	ATGAAGCCTA	AACTGATGTA	CCAGGAGCTG	AAGGTTCCTG	TTGAGGAACC	TGCGGGAGAA	CTGCCCATGA	70
5	ATGAAATCGA	GGCATGGAAG	GCAGCAGAGA	AGAAAGCCCG	TTGGGTCCTC	CTTGTCCTTA	TCCTGGCGGT	140
	AGTGGGCTTC	GGTGCCCTGA	TGACTCAGCT	GTTTCTATGG	GAATACGGGG	ACTTACATCT	ATTTGGCCCG	210
10	AATCAGCACC	CAGCCCCCTG	CTATGACCCC	TGCGAGGCGG	TGCTGGTGGA	GAGCATTCCC	GAGGGGCTGG	280
10	AGTTTCCCAA	TGCCACCACA	AGCAACCCCT	CCACCAGCCA	GGCCTGGTTG	GGCCTCCTTG	CCGGTGCTCA	350
	CAGCAGCCTG	GACATCGCGT	CCTTCTACTG	GACTCTCACA	AACAATGATA	CCCACACGCA	AGAGCCCTCT	420
15	GCCCAGCAGG	GTGAAGAGGT	TCTTCAGCAG	CTTCAGGCTC	TGGCACCTCG	AGGTGTAAAG	GTTCGCATCG	490
	CTGTGAGCAA	ACCCAACGGA	CCTCTGGCTG	ATCTGCAGTC	TCTGCTACAG	AGTGGTGCCC	AGGTGCGCAT	560
20	GGTGGACATG	CAGAAGCTGA	CCCATGGTGT	CCTGCACACC	AAGTTCTGGG	TGGTGGACCA	GACCCACTTT	630
20	TACCTGGGCA	GTGCCAACAT	GGACTGGCGA	TCGCTGACCC	AGGTCAAGGA	GCTGGGCGTG	GTCATGTACA	700
	ACTGCAGCTG	CCTGGCTCGC	GACCTCACCA	AGATTTTTGA	AGCCTATTGG	TTCCTGGGCC	AGGCAGGCAG	770
25	CTCCATCCCT	TCAACCTGGC	CACGGCCCTT	TGACACCCGG	TACAACCAAG	AAACACCGAT	GGAGATCTGC	840
	CTCAATGGCA	CCCCAGCCCT	GGCCTACCTG	GCGAGTGCAC	CCCCGCCACT	GTGTCCAGGT	GGCCGCACCC	910
30	CAGACCTGAA	GGCACTGCTC	AGCGTGGTGG	ACAACGCCCG	AAGCTTCATC	TACATTGCAG	TTATGAACTA	980
30	CCTGCCCACC	ATGGAGTTCT	CCCATCCACG	CAGGTTCTGG	CCAGCGATTG	ATGATGGGCT	AAGACGGGCT	1050
	GCGTATGAAC	GAGGCGTCAA	AGTGCGTTTG	CTCATCAGCT	GCTGGGGACA	CTCCGAGCCA	TCCATGCGGT	1120
35	CCTTCCTGCT	CTCCCTGGCT	GCCCTTCGTG	ACAACCATAC	CCACTCTGAC	ATCCAGGTGA	AACTGTTTGT	1190
	GGTCCCTGCG	GATGAGGCCC	AAGCTCGAAT	CCCCTATGCC	CGCGTCAACC	ACAACAAGTA	CATGGTGACT	1260
40	GAACGCACCA	CATACATTGG	AACCTCCAAC	TGGTCTGGAA	GCTACTTCAC	AGAGACGGCA	GGCACCTCCC	1330
40	TGCTGGTGAC	ACAGAACGGG	CACGGTGGCT	TGCGCAGCCA	GCTGGAGGCT	GTTTTCCTGA	GAGACTGGGA	1400
	ATCCCCATAC	AGCCACAACC	TTGACACCTC	AGCCGACAGT	GTGGGCAATG	CCTGCCGCCT	GCTTTGA 14	67

FIGURE 8

5

MKPKLMYQEL KVPVEEPAGE LPMNEIEAWK AAEKKARWVL LVLILAVVGF GALMTQLFLW EYGDLHLFGP 70

NQHPAPCYDP CEAVLVESIP EGLEFPNATT SNPSTSQAWL GLLAGAHSSL DIASFYWTLT NNDTHTQEPS 140

10 AQQGEEVLQQ LQALAPRGVK VRIAVSKPNG PLADLQSLLQ SGAQVRWVDM QKLTHGVLHT KFWVVDQTHF 210

YLGSANMDWR SLTQVKELGV VMYNCSCLAR DLTKIFEAYW FLGQAGSSIP STWPRPFDTR YNQETPMEIC 280

LNGTPALAYL ASAPPPLCPG GRTPDLKALL SVVDNARSFI YIAVMNYLPT MEFSHPRRFW PAIDDGLRRA 350

AYERGVKVRL LISCWGHSEP SMRSFLLSLA ALRDNHTHSD IQVKLFVVPA DEAQARIPYA RVNHNKYMVT 420

ERTTYIGTSN WSGSYFTETA GTSLLVTQNG HGGLRSQLEA VFLRDWESPY SHNLDTSADS VGNACRLL 488